

Refine Search

Search Results

Term	Documents
YEAST	186974
YEASTS	35004
(8 AND YEAST).PGPB,USPT,USOC,EPAB,JPAB,DWPI.	127
(L8 and yeast).PGPB,USPT,USOC,EPAB,JPAB,DWPI.	127

Database:

US Pre-Grant Publication Full-Text Database
 US Patents Full-Text Database
 US Patents OCR Backfile
 EPO Abstracts Database
 JPO Abstracts Database
 Derwent World Patents Index
 IBM Technical Disclosure Bulletin Database

Search:

L9

Refine Search

Recall Text

Clear

Interrupt

Search History

DATE: Monday, March 16, 2009 [Purge Queries](#) [Printable Copy](#) [Create Case](#)

<u>Set Name</u> side by side	<u>Query</u>	<u>Hit Count</u>	<u>Set Name</u> result set
<i>DB=PGPB,USPT,USOC,EPAB,JPAB,DWPI; THES=ASSIGNEE; PLUR=YES; OP=AND</i>			
<u>L9</u>	L8 and yeast	127	<u>L9</u>
<u>L8</u>	(katsuyama.in. or arakawa.in. or tokunaga.in. or yammamoto.in.) and (yeast)	127	<u>L8</u>
<u>L7</u>	L6 and yeast	0	<u>L7</u>
<u>L6</u>	changchun adj huapu.as.	22	<u>L6</u>
<u>L5</u>	L4 and respiration	69	<u>L5</u>
<u>L4</u>	(respiration adj yeast) or (deficient adj yeast)	1065	<u>L4</u>
<u>L3</u>	L2 and aspiration	2	<u>L3</u>
<u>L2</u>	L1 and respiration	153	<u>L2</u>

L1 (wang.in. or bao.in. or yu.in.) and yeast

4611 L1

END OF SEARCH HISTORY

Untitled

? e au=katsuyama, i?

Ref	Items	Index-term
E1	15	AU=KATSUYAMA, I
E2	17	AU=KATSUYAMA, I.
E3	0	AU=KATSUYAMA, I?
E4	12	AU=KATSUYAMA, ICHIRO
E5	1	AU=KATSUYAMA, ICHIROU
E6	316	AU=KATSUYAMA, ISAMU
E7	12	AU=KATSUYAMA, ITARU
E8	16	AU=KATSUYAMA, IWA0
E9	4	AU=KATSUYAMA, J
E10	30	AU=KATSUYAMA, J.
E11	16	AU=KATSUYAMA, JINYA
E12	1	AU=KATSUYAMA, JUICHI
E13	1	AU=KATSUYAMA, JUN'ICHI
E14	26	AU=KATSUYAMA, JUNJI
E15	12	AU=KATSUYAMA, JYUNJI
E16	11	AU=KATSUYAMA, K
E17	148	AU=KATSUYAMA, K.
E18	2	AU=KATSUYAMA, KANAKO
E19	1	AU=KATSUYAMA, KAORI
E20	9	AU=KATSUYAMA, KAYOKO
E21	2	AU=KATSUYAMA, KAZUHIKO
E22	27	AU=KATSUYAMA, KAZUKI
E23	12	AU=KATSUYAMA, KAZUO
E24	1	AU=KATSUYAMA, KEIICHI
E25	1	AU=KATSUYAMA, KEIKO

Enter PAGE for more

? s e8 and yeast

	16	AU=KATSUYAMA, IWA0
	1317956	YEAST
S1	4	AU='KATSUYAMA, IWA0' AND YEAST

? e au=arakawa, t?

Ref	Items	Index-term
E1	0	AU=ARAKAWA, T?
E2	32	AU=ARAKAWA, TADAHARU
E3	2	AU=ARAKAWA, TADAHIRO
E4	5	AU=ARAKAWA, TADAO
E5	7	AU=ARAKAWA, TADAOKI
E6	14	AU=ARAKAWA, TADASHI
E7	2	AU=ARAKAWA, TADAYOSHI
E8	1	AU=ARAKAWA, TADAYUKI
E9	6	AU=ARAKAWA, TAICHI
E10	1	AU=ARAKAWA, TAIJI
E11	42	AU=ARAKAWA, TAKAAKI
E12	3	AU=ARAKAWA, TAKACHIKA
E13	4	AU=ARAKAWA, TAKAFUMI
E14	4	AU=ARAKAWA, TAKAHARU
E15	23	AU=ARAKAWA, TAKAHIKO
E16	565	AU=ARAKAWA, TAKAHIRO
E17	2	AU=ARAKAWA, TAKAHISA
E18	26	AU=ARAKAWA, TAKAMASA
E19	22	AU=ARAKAWA, TAKAMI
E20	2	AU=ARAKAWA, TAKAMITSU
E21	2	AU=ARAKAWA, TAKANORI
E22	5	AU=ARAKAWA, TAKAO
E23	85	AU=ARAKAWA, TAKASHI
E24	11	AU=ARAKAWA, TAKATOSHI
E25	76	AU=ARAKAWA, TAKAYASU

Enter PAGE for more

Untitled

? e au=arakawa, tsutomu?

Ref	Items	Index-term
E1	0	AU=ARAKAWA, TSUTOMU?
E2	161	AU=ARAKAWA, TSUYOSHI
E3	1	AU=ARAKAWA, TSUYOSI
E4	2	AU=ARAKAWA, TUTOMU
E5	1	AU=ARAKAWA, TUYOSHI
E6	1	AU=ARAKAWA, U.
E7	1	AU=ARAKAWA, V.
E8	1	AU=ARAKAWA, W.
E9	3	AU=ARAKAWA, WATARU
E10	233	AU=ARAKAWA, Y
E11	2665	AU=ARAKAWA, Y.
E12	1	AU=ARAKAWA, Y. (RESEARCH CENTER FOR ADVANCED SCIE
E13	1	AU=ARAKAWA, Y. ET AL
E14	1	AU=ARAKAWA, Y. ET AL.
E15	1	AU=ARAKAWA, Y. H.
E16	32	AU=ARAKAWA, Y*
E17	1	AU=ARAKAWA, YASHICHIKA
E18	2	AU=ARAKAWA, YASHIHISA
E19	63	AU=ARAKAWA, YASUAKI
E20	915	AU=ARAKAWA, YASUHIKO
E21	20	AU=ARAKAWA, YASUHIRO
E22	3	AU=ARAKAWA, YASUKO
E23	2	AU=ARAKAWA, YASUMI
E24	9	AU=ARAKAWA, YASUNOBU
E25	2	AU=ARAKAWA, YASUNORI

Enter PAGE for more

? s e1-e2

	0	AU=ARAKAWA, TSUTOMU?
	161	AU=ARAKAWA, TSUYOSHI
S2	161	S E1-E2

? s s2 and yeast

	161	S2
	1317956	YEAST
S3	0	S S2 AND YEAST

? e au=masao, t?

Ref	Items	Index-term
E1	2	AU=MASAO, SUZUKI
E2	31	AU=MASAO, T.
E3	0	AU=MASAO, T?
E4	2	AU=MASAO, TANABE
E5	1	AU=MASAO, TANAKA
E6	1	AU=MASAO, TASAKA
E7	1	AU=MASAO, TATENO
E8	1	AU=MASAO, TOHRU
E9	1	AU=MASAO, TORU
E10	2	AU=MASAO, TOYODA
E11	1	AU=MASAO, TSURUOKA
E12	1	AU=MASAO, U.
E13	1	AU=MASAO, UEMURA
E14	3	AU=MASAO, USHIO
E15	4	AU=MASAO, Y.
E16	1	AU=MASAO, YAMADA
E17	1	AU=MASAO, YAMAZAKI
E18	1	AU=MASAO, YASUAKI YAMADA
E19	1	AU=MASAOA A P
E20	1	AU=MASAOA, A.
E21	3	AU=MASAOA, A. P.
E22	1	AU=MASAOA, P.

Untitled

E23 2 AU=MASAOAKA T
 E24 1 AU=MASAOAKA, T.
 E25 1 AU=MASAOCHIN
 Enter PAGE for more

? s eau=masao, tokunaga?

>>>w: One or more prefixes are unsupported
 or undefined in one or more files.

s4 0 S EAU=MASAO, TOKUNAGA?

? e au=tokunaga, masao?

Ref	Items	Index-term
E1	166	AU=TOKUNAGA, MASAO
E2	2	AU=TOKUNAGA, MASAO*
E3	0	AU=TOKUNAGA, MASAO?
E4	8	AU=TOKUNAGA, MASARU
E5	72	AU=TOKUNAGA, MASASHI
E6	2	AU=TOKUNAGA, MASASUKE
E7	3	AU=TOKUNAGA, MASATERU
E8	6	AU=TOKUNAGA, MASATO
E9	1	AU=TOKUNAGA, MASATOKI
E10	14	AU=TOKUNAGA, MASATOSHI
E11	2	AU=TOKUNAGA, MASATOSI
E12	8	AU=TOKUNAGA, MASAYA
E13	49	AU=TOKUNAGA, MASAYOSHI
E14	1	AU=TOKUNAGA, MASAYOSHI (KAGOSHIMA CITY HOSPITAL (
E15	3	AU=TOKUNAGA, MASAYUKI
E16	6	AU=TOKUNAGA, MASUNORI
E17	5	AU=TOKUNAGA, MAYUMI
E18	11	AU=TOKUNAGA, MICHIO
E19	4	AU=TOKUNAGA, MICHIO
E20	3	AU=TOKUNAGA, MIDORI
E21	3	AU=TOKUNAGA, MIE
E22	1	AU=TOKUNAGA, MIKI
E23	2	AU=TOKUNAGA, MIKIE
E24	6	AU=TOKUNAGA, MIKIKO
E25	1	AU=TOKUNAGA, MIKIO

Enter PAGE for more

? s e1-e3

	166	AU=TOKUNAGA, MASAO
	2	AU=TOKUNAGA, MASAO*
	0	AU=TOKUNAGA, MASAO?
s5	168	S E1-E3

? s s5 and yeast

	168	S5
	1317956	YEAST
s6	30	S S5 AND YEAST

? e au=yamamoto, tadashi?

Ref	Items	Index-term
E1	7	AU=YAMAMOTO, TADASHI S.
E2	1	AU=YAMAMOTO, TADASHI*
E3	0	AU=YAMAMOTO, TADASHI?
E4	2	AU=YAMAMOTO, TADASHIGE
E5	1	AU=YAMAMOTO, TADASHIO
E6	3	AU=YAMAMOTO, TADASI
E7	36	AU=YAMAMOTO, TADASU
E8	3	AU=YAMAMOTO, TADASUKE
E9	3	AU=YAMAMOTO, TADATERU
E10	21	AU=YAMAMOTO, TADATO
E11	127	AU=YAMAMOTO, TADATOSHI

Untitled

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E12      2  AU=YAMAMOTO, TADATSUGA
E13     47  AU=YAMAMOTO, TADATSUGA
E14     11  AU=YAMAMOTO, TADATSUKA
E15      5  AU=YAMAMOTO, TADAYASU
E16      6  AU=YAMAMOTO, TADAYOSHI
E17     22  AU=YAMAMOTO, TADAYUKI
E18     16  AU=YAMAMOTO, TAE
E19     26  AU=YAMAMOTO, TAEKA
E20      8  AU=YAMAMOTO, TAEKO
E21     29  AU=YAMAMOTO, TAHEI
E22      1  AU=YAMAMOTO, TAI
E23     44  AU=YAMAMOTO, TAIBO
E24      9  AU=YAMAMOTO, TAIBO (ED)
E25     77  AU=YAMAMOTO, TAICHI
```

Enter PAGE for more

? s e1-e2

```
      7  AU=YAMAMOTO, TADASHI S.
      1  AU=YAMAMOTO, TADASHI*
s7      8  S E1-E2
```

? s s7 and yeast

```
      8  S7
    1317956  YEAST
s8      0  S S7 AND YEAST
```

? s respiration and yeast

```
Processing
    1142014  RESPIRATION
    1317956  YEAST
s9      14717  S RESPIRATION AND YEAST
```

? s s9 and deficient

```
Processing
    14717  S9
    1239950  DEFICIENT
s10     2217  S S9 AND DEFICIENT
```

? s s10 and screen?

```
Processing
Processing
    2217  S10
    3391121  SCREEN?
s11     109  S S10 AND SCREEN?
```

? s s11 and respiration(w)deficient

```
    109  S11
    1142014  RESPIRATION
    1239950  DEFICIENT
    2386  RESPIRATION(W)DEFICIENT
s12     45  S S11 AND RESPIRATION(W)DEFICIENT
```

? rd

```
>>>w: Duplicate detection is not supported for File 393.
Duplicate detection is not supported for File 391.
Records from unsupported files will be retained in the RD set.
s13     13  RD (UNIQUE ITEMS)
```

? t s13/3,k/1-13

```
>>>w: KWIC option is not available in file(s): 399
13/3,K/1 (Item 1 from file: 5) Links
Fulltext available through: STIC Full Text Retrieval Options
Biosis Previews(R)
```

Untitled

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0020623649 Biosis No.: 200800670588

Intersection of RNA Processing and the Type II Fatty Acid Synthesis Pathway in Yeast Mitochondria

Author: Schonauer Melissa S; Kastaniotis Alexander J; Hiltunen J Kalervo; Dieckmann Carol L (Reprint)

Author Address: Univ Arizona, Dept Biochem and Mol Biophys, POB 210106, Tucson, AZ 85721 USA**USA

Author E-mail Address: dieckman@u.arizona.edu

Journal: Molecular and Cellular Biology 28 (21): p 6646-6657 NOV 2008 2008

Item Identifier: doi:10.1128/MCB.01162-08

ISSN: 0270-7306

Document Type: Article

Record Type: Abstract

Language: English

Intersection of RNA Processing and the Type II Fatty Acid Synthesis Pathway in Yeast Mitochondria

Abstract: Distinct metabolic pathways can intersect in ways that allow hierarchical or reciprocal regulation. In a screen of respiration- deficient *Saccharomyces cerevisiae* gene deletion strains for defects in mitochondrial RNA processing, we found that lack... ..pathway is the sole precursor for lipoic acid synthesis and attachment. The protein component of yeast mitochondrial RNase P, Rpm2, is not modified by lipoic acid in the wild-type strain...

13/3,K/2 (Item 2 from file: 5) Links

Fulltext available through: STIC Full Text Retrieval Options

Biosis Previews(R)

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17372760 Biosis No.: 200300331056

Toxicity of metal ions used in dental alloys: A study in the yeast *Saccharomyces cerevisiae*.

Author: Yang Hyeong-Cheol; Pon Liza A (Reprint)

Author Address: Department of Anatomy and Cell Biology, Columbia University, 630 West 168th St., P and S 12-425, New York, NY, 10032, USA**USA

Author E-mail Address: lap5@columbia.edu

Journal: Drug and Chemical Toxicology an International Journal for Rapid Communication 26 (2): p 75-85 May 2003 2003

Medium: print

ISSN: 0148-0545

Document Type: Article

Record Type: Abstract

Language: English

Toxicity of metal ions used in dental alloys: A study in the yeast *Saccharomyces cerevisiae*.

Abstract: ...In this study, we investigated the effect of metal ions on growth of the budding yeast, *Saccharomyces cerevisiae*, and on the morphology and function of yeast mitochondria. Moreover, we tested whether mitochondrial respiratory activity contributes to metal toxicity. Metal ions affected yeast cell growth. The toxicity of metal ions to yeast cells, ranked in decreasing order are as follows: Hg > Ag > Au > Cu, Ni, Co, Zn... ..respectively. None of the toxic metal ions resulted in loss of mitochondrial respiratory activity. However, respiration- deficient rho0 cells appeared to be resistant to Ag ion, but not to Hg and Au... ..Furthermore, at high concentrations, Ag ion caused morphological changes in mitochondria. These studies indicate that yeast may be used as a model system to screen for toxic effect of metals ions from dental alloys, and that oxidation activity in mitochondria...

Untitled

13/3,K/3 (Item 3 from file: 5) Links
Fulltext available through: STIC Full Text Retrieval Options
Biosis Previews(R)
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16961532 Biosis No.: 200200555043
Merging mitochondria matters. Cellular role and molecular machinery of mitochondrial fusion

Author: Westermann Benedikt (Reprint)
Author Address: Institut fuer Physiologische Chemie der Universitaet Muenchen,
Butenandtstrasse 5, D-81377, Muenchen, Germany**Germany
Journal: EMBO Reports 3 (6): p 527-531 June, 2002 2002
Medium: print
ISSN: 1469-221X
Document Type: Article; Literature Review
Record Type: Abstract
Language: English

Abstract: Fusion is essential for mitochondrial function in a great variety of eukaryotic cell types. Yeast cells defective in mitochondrial fusion are respiration-deficient, human cells use complementation of fused mitochondria as a defence against the accumulation of oxidative... ..evolutionarily conserved large GTPase in the outer membrane is essential for mitochondrial fusion, and genetic screens in yeast are revealing an increasing number of additional important genes. Mechanistic studies have provided the first...

DESCRIPTORS:
Organisms: ...yeast (Fungi
Organisms: Parts Etc:
Methods & Equipment: genetic screening--
Geographical Name:

13/3,K/4 (Item 4 from file: 5) Links
Fulltext available through: STIC Full Text Retrieval Options
Biosis Previews(R)
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15804067 Biosis No.: 200000522380
No mutagenic or recombinogenic effects of mobile phone fields at 900 MHz detected in the yeast *Saccharomyces cerevisiae*

Author: Gos Pascal; Eicher Bernhard; Kohli Jurg; Heyer Wolf-Dietrich (Reprint)
Author Address: Section of Microbiology, University of California, Davis, One Shields Avenue, Davis, CA, 95616, USA**USA
Journal: Bioelectromagnetics 21 (7): p 515-523 October, 2000 2000
Medium: print
ISSN: 0197-8462
Document Type: Article
Record Type: Abstract
Language: English
No mutagenic or recombinogenic effects of mobile phone fields at 900 MHz detected in the yeast *Saccharomyces cerevisiae*

Abstract: Both actively growing and resting cells of the yeast *Saccharomyces cerevisiae* were exposed to 900-MHz fields that closely matched the Global System for... ..specific forward mutation assay at CAN1 and a wide-range assay measuring the induction of respiration-deficient (petite) clones that have lost their mitochondrial function. In addition, two further assays measured the... ..possible effects on genomic stability: First, an intrachromosomal, deletion-formation assay previously developed for genotoxic screening; and second, an intragenic recombination assay in the ADE2 gene. Fluctuation tests failed to detect...

DESCRIPTORS:

Untitled

Methods & Equipment: ...genotoxic screening--... ...screening method
Geographical Name:
Miscellaneous Terms: Concept Codes: ...respiration;yeast mutation rates

13/3,K/5 (Item 5 from file: 5) Links

Fulltext available through: STIC Full Text Retrieval Options
Biosis Previews(R)

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11280242 Biosis No.: 199293123133

STRUCTURE AND FUNCTION OF MRP20 AND MRP49 THE NUCLEAR GENES FOR TWO PROTEINS OF THE 54S SUBUNIT OF THE YEAST MITOCHONDRIAL RIBOSOME

Author: FEARON K (Reprint); MASON T L

Author Address: DEP BIOCHEM, PROGRAM MOLECULAR CELLULAR BIOL, UNIV MASSACHUSETTS, AMHERST, MASSACHUSETTS 01003, USA**USA

Journal: Journal of Biological Chemistry 267 (8): p 5162-5170 1992

ISSN: 0021-9258

Document Type: Article

Record Type: Abstract

Language: ENGLISH

...MRP20 AND MRP49 THE NUCLEAR GENES FOR TWO PROTEINS OF THE 54S SUBUNIT OF THE YEAST MITOCHONDRIAL RIBOSOME

Abstract: ...large subunit of the mitochondrial ribosome in *Saccharomyces cerevisiae*. Their genes were identified through immunological screening of a genomic library in the expression vector λ gt11. Nucleotide sequencing revealed that MRP49... ...of mutations in essential genes for mitochondrial translation. Inactivation of MRP49 caused a cold-sensitive respiration- deficient phenotype, indicating that MRP49 is not an essential ribosomal protein. The *mrp49* mutants were defective... ...the nonpermissive temperature. With the results presented here, there are now published sequences for 14 yeast mitochondrial ribosomal proteins, only five of which bear discernable relationships to eubacterial ribosomal proteins.

13/3,K/6 (Item 1 from file: 24) Links

Fulltext available through: STIC Full Text Retrieval Options
CSA Life Sciences Abstracts

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0002323543 IP Accession No: 5379117

The mitochondrial cytochrome c peroxidase *Ccp1* of *Saccharomyces cerevisiae* is involved in conveying an oxidative stress signal to the transcription factor *Pos9* (*Skn7*)

Charizanis, C; Juhnke, H; Krems, B; Entian, K-D Institut für Mikrobiologie, Johann Wolfgang Goethe-Universität Frankfurt, Biozentrum, Niederursel, Marie-Curie-Strasse 9, D-60439 Frankfurt am Main, Germany, [mailto:krems@em.uni-frankfurt.de]

Molecular and General Genetics , v 262 , n 3 , p 437-447 , 1999

Publication Date: 1999

Publisher: Springer-Verlag (Berlin), Heidelberger Platz 3 Berlin 14197 Germany, [mailto:subscriptions@springer.de]

Document Type: Journal Article

Record Type: Abstract

Language: English

Summary Language: English

ISSN: 0026-8925

Electronic ISSN: 1432-1874; 1432-1874

File Segment: Nucleic Acids Abstracts; Algology, Mycology & Protozoology Abstracts (Microbiology C)

Abstract:

Untitled

...which are involved in the oxygen-dependent activation of the Gal4-Pos9 hybrid protein we screened for mutants that failed to induce the heterologous test system upon oxidative stress (fap mutants for factors activating Pos9). We isolated several respiration-deficient and some respiration-competent mutants by this means. We selected for further characterization only those mutants which also displayed an oxidative-stress-sensitive phenotype. One of the respiration-deficient mutants (complementation group fap6) could be complemented by the ISM1 gene, which encodes mitochondrial isoleucyl tRNA synthetase, suggesting that respiration competence was important for signalling of oxidative stress. In accordance with this notion a rho0 strain and a wild-type strain in which respiration had been blocked (by treatment with antimycin A or with cyanide) also failed to activate Gal4-Pos9 upon imposition of oxidative stress. Another mutant, fap24, which was respiration-competent, could be complemented by CCP1, which encodes the mitochondrial cytochrome c peroxidase. Mitochondrial cytochrome...

Descriptors: Oxidative stress; Cyanide; Isoleucine-tRNA ligase; Transcription factors; Complementation; Reporter gene; Respiration; Mitochondria; Pos9 protein; Ccp1 protein; ISM1 gene; antimycin A; Skn7 protein; cytochrome-c peroxidase; Saccharomyces...
Identifiers: budding yeast

13/3,K/7 (Item 1 from file: 34) Links

Fulltext available through: STIC Full Text Retrieval Options
SciSearch(R) Cited Ref Sci

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13589951 Genuine Article#: 895PD No. References: 86

The importance of mutation, then and now: studies with yeast cytochrome c

Author: Sherman F (REPRINT)

Corporate Source: Univ Rochester, Sch Med, Dept Biochem & Biophys, Box 712/Rochester//NY/14642 (REPRINT); Univ Rochester, Sch Med, Dept Biochem & Biophys, Rochester//NY/14642 (fred.sherman@urmc.rochester.edu)

Journal: MUTATION RESEARCH-REVIEWS IN MUTATION RESEARCH, 2005, V 589, N1 (JAN), P 1-16

ISSN: 1383-5742 Publication date: 20050100

Publisher: ELSEVIER SCIENCE BV, PO BOX 211, 1000 AE AMSTERDAM, NETHERLANDS

Language: English Document Type: ARTICLE (ABSTRACT AVAILABLE)

The importance of mutation, then and now: studies with yeast cytochrome c

Abstract: ...initiation mutants, allowing the early cloning of the gene. A method was developed for transforming yeast directly with synthetic oligonucleotides, resulting in the convenient production of CYC1 mutants with defined sequences... nucleus, and on N-terminal acetylation stems from properties of CYC1 mutants isolated in early screens more than a decade ago. (C) 2004 Elsevier B.V. All rights reserved.

Identifiers-- ...AMINO-ACID REPLACEMENTS; ALPHA-TERMINAL ACETYLTRANSFERASE; RESPIRATION-DEFICIENT MUTANTS; CYC1 MESSENGER-RNA; SACCHAROMYCES-CEREVISIAE; SYNTHETIC OLIGONUCLEOTIDES; ISO-1-CYTOCHROME C; BAKERS-YEAST; IN-VIVO; METHIONINE AMINOPEPTIDASE

13/3,K/8 (Item 2 from file: 34) Links

Fulltext available through: STIC Full Text Retrieval Options
SciSearch(R) Cited Ref Sci

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13387935 Genuine Article#: 874ZN No. References: 126

Mitochondrial genomics and proteomics

Author: Schonauer MS; Dieckmann CL (REPRINT)

Corporate Source: Univ Arizona, Dept Biochem & Mol Biophys, Tucson//AZ/85721 (REPRINT); Univ Arizona, Dept Biochem & Mol Biophys, Tucson//AZ/85721; Univ Arizona, Dept Mol & Cellular Biol, Tucson//AZ/85721 (dieckman@u.arizona.edu)

Untitled

Journal: CURRENT GENOMICS , 2004 , V 5 , N7 (NOV) , P 575-588
ISSN: 1389-2029 Publication date: 20041100
Publisher: BENTHAM SCIENCE PUBL LTD , EXECUTIVE STE Y26, PO BOX 7917, SAIF ZONE,
1200 BR SHARJAH, U ARAB EMIRATES
Language: English Document Type: REVIEW (ABSTRACT AVAILABLE)
Abstract: Factors involved in mitochondrial biogenesis and function have been
studied classically via mutagenesis screens and subsequent genetic and biochemical
analyses. The recent advent of high-throughput technologies has provided... ..will
describe past and present genomic and proteomic methods used to study mitochondria
both in yeast and mammalian cells, their advantages and limitations, and the current
knowledge of the number of...
Identifiers-- ...RESPIRATION-DEFICIENT MUTANTS; CONSERVED DODECAMER SEQUENCE;
CYTOCHROME-C-OXIDASE; SACCHAROMYCES-CEREVISIAE; GENE-EXPRESSION; MESSENGER-RNA;
NUCLEAR GENE

13/3,K/9 (Item 3 from file: 34) Links
SciSearch(R) Cited Ref Sci
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01557284 Genuine Article#: HH747 No. References: 58
STRUCTURE AND FUNCTION OF MRP20 AND MRP49, THE NUCLEAR GENES FOR 2 PROTEINS OF THE
54-S-SUBUNIT OF THE YEAST MITOCHONDRIAL RIBOSOME

Author: FEARON K; MASON TL
Corporate Source: UNIV MASSACHUSETTS,DEPT BIOCHEM/AMHERST//MA/01003; UNIV
MASSACHUSETTS,DEPT BIOCHEM/AMHERST//MA/01003; UNIV MASSACHUSETTS,PROGRAM MOLEC &
CELLULAR BIOL/AMHERST//MA/01003
Journal: JOURNAL OF BIOLOGICAL CHEMISTRY , 1992 , V 267 , N8 (MAR 15) , P
5162-5170
Language: ENGLISH Document Type: ARTICLE (Abstract Available)
...AND MRP49, THE NUCLEAR GENES FOR 2 PROTEINS OF THE 54-S-SUBUNIT OF THE YEAST
MITOCHONDRIAL RIBOSOME

Abstract: ...large subunit of the mitochondrial ribosome in saccharomyces
cerevisiae. Their genes were identified through immunological screening of a genomic
library in the expression vector lambda-gt11. Nucleotide sequencing revealed that
MRP49... ..of mutations in essential genes for mitochondrial translation.
Inactivation of MRP49 caused a cold-sensitive respiration- deficient phenotype,
indicating that MRP49 is not an essential ribosomal protein. The mrp49 mutants were
defective... ..the nonpermissive temperature. With the results presented here,
there are now published sequences for 14 yeast mitochondrial ribosomal proteins,
only five of which bear discernable relationships to eubacterial ribosomal proteins.

Identifiers--
Research Fronts: ...FASCIANS CLONING VECTORS; ESCHERICHIA-COLI CHROMOSOME; PRECISE
IDENTIFICATION)
90-2893 002 (GENE TARGETING; HOMOLOGOUS RECOMBINATION; YEAST NUCLEAR-FUSION;
EMBRYONIC STEM-CELLS)
90-1410 001 (MITOCHONDRIAL PROTEIN IMPORT; DEFICIENCY IN 2 YEAST COENZYME-Q
MUTANTS; STRUCTURAL GENE ENCODING HEXAPRENYL PYROPHOSPHATE SYNTHETASE)
90-4609 001 (5S RIBOSOMAL-RNA...
Cited References:

13/3,K/10 (Item 1 from file: 73) Links
Fulltext available through: STIC Full Text Retrieval Options
EMBASE
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0070167499 EMBASE No: 1974169087
A synthesis of labeled ethidium bromide

De Nobrega Bastos R.; Mahler H.R.
Dept. Chem., Indiana Univ., Bloomington, Ind. 47401, United States

Untitled

Corresp. Author/Affil: : Dept. Chem., Indiana Univ., Bloomington, Ind. 47401, United States

Archives of Biochemistry and Biophysics (ARCH. BIOCHEM. BIOPHYS.) January 1, 1974 , 160/2 (643-646)

CODEN: ABBIA ISSN: 0003-9861

Document Type: Journal ; Article Record Type: Abstract

Language: English

...its absorption spectra (uv, visible, ir), chromatographic behavior, and mutagenic effectiveness in the induction of respiration deficient cell lines in baker's yeast.

Medical Descriptors:

* drug screening; *drug synthesis; *pharmacology

13/3,K/11 (Item 1 from file: 370) Links

Science

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00509183 (USE 9 FOR FULLTEXT)

The Machinery of Mitochondrial Inheritance and Behavior

Yaffe, Michael P.<CRF RID="C1">

The author is in the Department of Biology, University of California, San Diego, La Jolla, CA 92093-0347, USA.

Science Vol. 283 5407 pp. 1493

Publication Date: 3-05-1999 (990305) Publication Year: 1999

Document Type: Journal ISSN: 0036-8075

Language: English

Section Heading: REVIEWS

Word Count: 4200 (THIS IS THE FULLTEXT)

Text:

...approaches. The first molecules mediating mitochondrial inheritance were discovered through isolation and analysis of mutant yeast cells that displayed conditional defects in mitochondrial distribution and morphology (B13) . More recently, proteins facilitating...

...types including mammalian neurons, sperm cells, and cultured fibroblasts as well as in the fission yeast *Schizosaccharomyces pombe* and the protozoan *Acanthamoeba castellanii* (B18) (B24) (B25) (B26) . Involvement of microtubules was...microtubule drugs, mitochondrial movement in neuronal axons became dependent on actin filaments (B29) . In the yeast *Saccharomyces cerevisiae*, a fraction of the mitochondrial tubules appears to colocalize with actin cables (B40)...

...mutations in the single actin gene cause abnormal mitochondrial distribution and morphology (B40) . Additionally, isolated yeast mitochondria bound to actin filaments and exhibited actin-based motility in an in vitro assay...

...these actin cables, but mitochondrial transport remains normal (B40) (B43) . Furthermore, mutations in the five yeast myosin genes have no effect on mitochondrial inheritance (B42) (B44) . Further clarification of actin's...to DNML1, a gene encoding a dynamin-like protein most closely related to Drp1 (B47) . Yeast cells expressing Dnm1p mutated in the conserved GTP-binding site also display collapsed mitochondria. Wild...

...protein sequence provides few hints as to its molecular function, but the similarity of the yeast mutant phenotype to that caused by mutations in dnml suggests that these two proteins may...insights into the function of the fuzzy onions protein have emerged from studies of the

Untitled

yeast homolog, Fzo1p (B52) (B53) . Mutations in Fzo1p caused fragmentation of mitochondrial tubules and eventual loss of mitochondrial DNA. Mitochondrial fusion, which normally follows the mating of two yeast cells, did not occur when Fzo1p was defective or absent, providing direct evidence of a...

...Mitochondrial Inheritance Mutants In contrast to animal cells, the yeast *S. cerevisiae* proliferates by a budding process in which a mother cell produces a daughter...

...of mitochondria and other organelles into the developing bud (B55) (Fig. 2). The analysis of yeast mutants defective in this transport process has led to the identification of proteins that facilitate...

...The first mutants affecting mitochondrial inheritance were isolated by microscopically screening collections of temperature-sensitive strains for cells that failed to transport mitochondria into daughter buds ...

...mitochondrial shape, and the mutants were named mdm for mitochondrial distribution and morphology. Although other yeast mutants that lose mitochondrial DNA and become respiration-deficient still retain mitochondrial compartments and remain viable (B56) , buds of mdm mutants devoid of mitochondria...

...in vitro into 10-nm-diameter filaments closely resembling IFs in animal cells (B59) . In yeast cells, Mdm1p localizes to punctate structures throughout the cytoplasm, and under conditions leading to the...classical ultrastructural features including double membranes and inner membrane cristae and remain partially competent for respiration, but are defective for division and inheritance by daughter buds. Mdm10p, Mmm1p, and Mdm12p are...

...mitochondrial distribution and morphology. Protein phosphorylation may also regulate mitochondrial inheritance, as mitochondrial transmission to yeast buds is delayed by mutations in the gene encoding serine-threonine phosphatase Ptc1p (B71...KIF5B, Mouse; fly Microtubules/ Mitochondrial aggregation near

KIF1B;			mitochondrial nucleus
KLP67A			surface
Drp1; Dnm1p	Human; yeast	Cytoplasm	Mitochondrial aggregation, reduced lateral distribution and/or branching
CluA; Clu1p	Slime mold; yeast	Cytoplasm	Mitochondrial aggregation, reduced lateral distribution and/or branching
Mdm1p	Yeast	Cytoplasm	Defective mitochondrial transmission to buds; fragmentation of tubules
Mdm20p	Yeast	Cytoplasm	Defective mitochondrial transmission to buds
Mdm14p	Yeast	Cytoplasm	Defective mitochondrial transmission to buds; mitochondrial aggregation
Rsp5p	Yeast	Cytoplasm	Defective mitochondrial transmission to buds; mitochondrial aggregation
Mdm10p	Yeast; fungus	Mitochondrial outer membrane	Defective mitochondrial transmission to buds; giant spherical mitochondria
Mmm1p	Yeast	Mitochondrial outer membrane	Defective mitochondrial transmission to buds; giant spherical mitochondria
Mdm12p	Yeast	Mitochondrial	Defective mitochondrial

			Untitled
Mgm1p	Yeast	outer membrane Mitochondrial	transmission to buds; giant spherical mitochondria Defective mitochondrial
Fuzzy fusion; onions, Fz0lp	Fly; yeast	outer membrane Mitochondrial	transmission to buds; mitochondrial aggregation Aberrant mitochondrial
End Table: Columns 1...		outer membrane	fragmentation of tubules

...Table: Columns 5 - 6 of 6 tblend

Figure F2

Caption: Mitochondrial network in the budding yeast *S. cerevisiae*. Wild-type yeast cells were engineered to express green fluorescent protein fused to a mitochondrial targeting sequence (the...

References and Notes:

...by Strathern, J. N., Jones, E. W., Broach, J. R., The Molecular Biology of the Yeast *Saccharomyces cerevisiae*, vol. 1 1981, 471504 Cold Spring Harbor Laboratory Press, Cold Spring Harbor, NY...

...TEXT:

Saccharomyces cerevisiae gene deletion strains for defects in mitochondrial RNA processing, we found that lack of any enzyme in the mitochondrial fatty acid type II biosynthetic pathway (FAS II) led to inefficient 5' processing...

...
Rpm2, is not modified by lipoic acid in the wild-type strain, and it is imported in FAS II mutant strains. Thus, a product of the FAS II pathway is required...

...
Cellular Biology (Intersection of RNA Processing and the Type II Fatty Acid Synthesis Pathway in Yeast Mitochondria. *Molecular and Cellular Biology*, 2008;28(21):6646-6657). For more information, contact C...

13/3,K/13 (Item 1 from file: 357) Links

Derwent Biotech Res.

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0306690 DBA Accession No.: 2003-08475 PATENT

Identifying secondary target site in telomerase pathway, in which mutations in the secondary target site provide unviable host cells when the telomerase gene is overexpressed, useful for screening drugs for cancer expressed sequence tag and transgenic mouse for use in gene therapy and drug screening

Patent Assignee: HUTCHINSON CANCER RES CENT FRED 2002

Patent Number: WO 200290588 Patent Date: 20021114 WPI Accession No.: 2003-111990

(200310)

Priority Application Number: WO 200027430 Application Date: 20001003

National Application Number: WO 2000US27430 Application Date: 20001003

Language: English

...secondary target site provide unviable host cells when the telomerase gene is overexpressed, useful for screening drugs for cancer expressed sequence tag and transgenic mouse for use in gene therapy and drug screening

Untitled

Abstract: ...downregulation, elimination or disruption of (II) gives rise to senescence or synthetic lethality; and (6) screening for drugs involves providing one or more eukaryotic cells capable of telomerase overexpression, which one... precursor, AAD10 or their combinations. (M1) further involves using (I), or its lethal mutations, to screen for a drug or drug candidate which comprises a polypeptide, oligonucleotide, polysaccharide, or small molecule... body weight/day. EXAMPLE - Mutants which were inviable in the presence of telomerase overexpression in yeast or vice versa were screened. The yeast strain used contained the expressed sequence tag (EST)1, EST2 and TEL1 genes under the...containing media, but grew on dextrose and glycerol-containing media were selected. Colonies formed by respiration-deficient cells did not grow on glycerol and thus were omitted from further analysis. The isolates...

E.C. Numbers:

Descriptors: telomerase pathway secondary target site mutation det., drug screening, transgenic mouse, EST1, EST2, EST3, TLC1, embZ35904, gbU14595, embZ35905, dbjD28120, gbL24113, embX76992, gbAC005476.3, gbU53340 ...